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1600

RAW SEQUENCE LISTING

PATENT APPLICATION: US/09/905,114A

DATE: 03/25/2003

TIME: 12:54:15

Input Set : A:\210707US.txt

Output Set: N:\CRF4\03252003\I905114A.raw

3 <110> APPLICANT: ZHANG, HUANMIN
 4 AX, ROY L
 5 BELLIN, MARY E
 7 <120> TITLE OF INVENTION: ISOLATED POLYNUCLEOTIDE SEQUENCES ENCODING A FERTILITY
 ASSOCIATED ANTIGEN
 9 <130> FILE REFERENCE: 210707US20
 11 <140> CURRENT APPLICATION NUMBER: 09/905,114A
 12 <141> CURRENT FILING DATE: 2001-07-16
 14 <150> PRIOR APPLICATION NUMBER: US 60/218,140
 15 <151> PRIOR FILING DATE: 2000-07-14
 17 <160> NUMBER OF SEQ ID NOS: 10
 19 <170> SOFTWARE: PatentIn version 3.1
 21 <210> SEQ ID NO: 1
 22 <211> LENGTH: 592
 23 <212> TYPE: DNA
 24 <213> ORGANISM: Bos sp.
 26 <220> FEATURE:
 27 <221> NAME/KEY: CDS
 28 <222> LOCATION: (1)..(591)
 29 <223> OTHER INFORMATION:
 W--> 31 <400> 1

ENTERED

32	gag aag cta aac gga aat tca aga aaa ggc ata aca tac aac tat gtg	48
33	Glu Lys Leu Asn Gly Asn Ser Arg Lys Gly Ile Thr Tyr Asn Tyr Val	
34	1 5 10 15	
36	att agc tct cgc ctt gga aga aac aca tat aaa gaa cag tat gcc ttt	96
37	Ile Ser Ser Arg Leu Gly Arg Asn Thr Tyr Lys Glu Gln Tyr Ala Phe	
38	20 25 30	
40	ctc tat aaa gaa aag cta gtg tct gta aaa caa agc tac ctc tac cac	144
41	Leu Tyr Lys Glu Lys Leu Val Ser Val Lys Gln Ser Tyr Leu Tyr His	
42	35 40 45	
44	gac tat cag gct gga gac gca gat gtg ttt tcc agg gaa ccc ttt gtg	192
45	Asp Tyr Gln Ala Gly Asp Ala Asp Val Phe Ser Arg Glu Pro Phe Val	
46	50 55 60	
48	gtc tgg ttc cag tca ccc tac acc gct gtc aag gac ttc gtg att gtc	240
49	Val Trp Phe Gln Ser Pro Tyr Thr Ala Val Lys Asp Phe Val Ile Val	
50	65 70 75 80	
52	ccc ctg cac acc acc cct gag aca tcc gtt aga gag att gat gag ctg	288
53	Pro Leu His Thr Thr Pro Glu Thr Ser Val Arg Glu Ile Asp Glu Leu	
54	85 90 95	
56	gct gat gtc tac aca gat gtg aaa cgt cgc tgg aat gca gag aat ttc	336
57	Ala Asp Val Tyr Thr Asp Val Lys Arg Arg Trp Asn Ala Glu Asn Phe	
58	100 105 110	
60	att ttc atg ggt gac ttc aat gct ggc tgc agc tac gtc ccc aag aag	384
61	Ile Phe Met Gly Asp Phe Asn Ala Gly Cys Ser Tyr Val Pro Lys Lys	

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62	115	120	125	
64	gcc tgg aag gac atc cgc ctg agg acg gac ccc aag ttc gtt tgg ctg			432
65	Ala Trp Lys Asp Ile Arg Leu Arg Thr Asp Pro Lys Phe Val Trp Leu			
66	130	135	140	
68	atc ggg gac caa gag gac acc acg gtc aag aag agc aca aac tgc gcc			480
69	Ile Gly Asp Gln Glu Asp Thr Thr Val Lys Lys Ser Thr Asn Cys Ala			
70	145	150	155	160
72	tat gac agg atc gtg ctt aga gga caa aat att gtc aac tct ggt ggt			528
73	Tyr Asp Arg Ile Val Leu Arg Gly Gln Asn Ile Val Asn Ser Gly Gly			
74	165	170	175	
76	cct caa tca aac ctc gtc ttt gat ttc cag aaa gct tac agg ttg tct			576
77	Pro Gln Ser Asn Leu Val Phe Asp Phe Gln Lys Ala Tyr Arg Leu Ser			
78	180	185	190	
80	gaa tcg aag gcc ctg g			592
81	Glu Ser Lys Ala Leu			
82	195			
85	<210> SEQ ID NO: 2			
86	<211> LENGTH: 197			
87	<212> TYPE: PRT			
88	<213> ORGANISM: Bos sp.			
90	<400> SEQUENCE: 2			
92	Glu Lys Leu Asn Gly Asn Ser Arg Lys Gly Ile Thr Tyr Asn Tyr Val			
93	1	5	10	15
96	Ile Ser Ser Arg Leu Gly Arg Asn Thr Tyr Lys Glu Gln Tyr Ala Phe			
97	20	25	30	
100	Leu Tyr Lys Glu Lys Leu Val Ser Val Lys Gln Ser Tyr Leu Tyr His			
101	35	40	45	
104	Asp Tyr Gln Ala Gly Asp Ala Asp Val Phe Ser Arg Glu Pro Phe Val			
105	50	55	60	
108	Val Trp Phe Gln Ser Pro Tyr Thr Ala Val Lys Asp Phe Val Ile Val			
109	65	70	75	80
112	Pro Leu His Thr Thr Pro Glu Thr Ser Val Arg Glu Ile Asp Glu Leu			
113	85	90	95	
116	Ala Asp Val Tyr Thr Asp Val Lys Arg Arg Trp Asn Ala Glu Asn Phe			
117	100	105	110	
120	Ile Phe Met Gly Asp Phe Asn Ala Gly Cys Ser Tyr Val Pro Lys Lys			
121	115	120	125	
124	Ala Trp Lys Asp Ile Arg Leu Arg Thr Asp Pro Lys Phe Val Trp Leu			
125	130	135	140	
128	Ile Gly Asp Gln Glu Asp Thr Thr Val Lys Lys Ser Thr Asn Cys Ala			
129	145	150	155	160
132	Tyr Asp Arg Ile Val Leu Arg Gly Gln Asn Ile Val Asn Ser Gly Gly			
133	165	170	175	
136	Pro Gln Ser Asn Leu Val Phe Asp Phe Gln Lys Ala Tyr Arg Leu Ser			
137	180	185	190	
140	Glu Ser Lys Ala Leu			
141	195			
144	<210> SEQ ID NO: 3			
145	<211> LENGTH: 671			

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146 <212> TYPE: DNA
 147 <213> ORGANISM: Bos sp.
 149 <400> SEQUENCE: 3
 150 acaacaggat ctgccccata ctgatggaga agctaaacgg aaattcaaga aaaggcataa 60
 152 catacaacta tgtgattagc tctcgccctg gaagaaacac atataaagaa cagttatgcct 120
 154 ttctctataa agaaaagcta gtgtctgtaa aacaaagcta cctctaccac gactatcagg 180
 156 ctggagacgc agatgtgttt tccagggAAC cctttgtggc ctggttccag tcaccctaca 240
 158 ccgctgtcaa ggacttcgtg attgtcccccc tgacacaccac ccctgagaca tccgttagag 300
 160 agattgtatgat gctggctgat gtctacacag atgtgaaacg tcgctggat gcagagaatt 360
 162 tcattttcat ggggtgacttc aatgctggct gcagctacgt ccccaagaag gcctggagg 420
 164 acatccgcct gaggacggac cccaaagttcg tttggctgat cggggaccaa gaggacacca 480
 166 cggtaagaa gggcacaaac tgccctatg acaggatcg tcttagagga caaaatattg 540
 168 tcaactctgg tggtccctcaa tcaaaccctcg tctttgattt ccagaaagct tacagggtgt 600
 170 ctgaatcgaa ggccctggat gtcagcgacc acttccagt tcatacatcat catcatcatg 660
 172 aagaaccatg a 671
 175 <210> SEQ ID NO: 4
 176 <211> LENGTH: 21
 177 <212> TYPE: DNA
 178 <213> ORGANISM: Artificial Sequence
 180 <220> FEATURE:
 181 <223> OTHER INFORMATION: synthetic DNA
 183 <400> SEQUENCE: 4
 184 cgtgaggagc ttccggcgaga g 21
 187 <210> SEQ ID NO: 5
 188 <211> LENGTH: 26
 189 <212> TYPE: PRT
 190 <213> ORGANISM: Bos sp.
 192 <220> FEATURE:
 193 <221> NAME/KEY: misc_feature
 194 <222> LOCATION: (4)..(4) /
 195 <223> OTHER INFORMATION: Xaa is any amino acid
 198 <400> SEQUENCE: 5
 W--> 200 Leu Lys Ile Xaa Ser Phe Asn Val Arg Ser Phe Gly Glu Ser Lys Lys
 201 1 5 10 15
 204 Ala Gly Phe Asn Ala Met Arg Val Ile Val
 205 20 25
 208 <210> SEQ ID NO: 6
 209 <211> LENGTH: 26
 210 <212> TYPE: DNA
 211 <213> ORGANISM: Artificial Sequence
 213 <220> FEATURE:
 214 <223> OTHER INFORMATION: synthetic DNA
 216 <400> SEQUENCE: 6
 217 acaacaggat ctgccccata ctgatg 26
 220 <210> SEQ ID NO: 7
 221 <211> LENGTH: 25
 222 <212> TYPE: DNA
 223 <213> ORGANISM: Artificial Sequence
 225 <220> FEATURE:

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226 <223> OTHER INFORMATION: synthetic DNA
228 <400> SEQUENCE: 7
229 tcaactggaa agtggtcgct gacat 25
232 <210> SEQ ID NO: 8
233 <211> LENGTH: 27
234 <212> TYPE: DNA
235 <213> ORGANISM: Artificial Sequence
237 <220> FEATURE:
238 <223> OTHER INFORMATION: synthetic DNA
240 <400> SEQUENCE: 8
241 aacaacaggat ctgccccata ctgatgg 27
244 <210> SEQ ID NO: 9
245 <211> LENGTH: 57
246 <212> TYPE: DNA
247 <213> ORGANISM: Artificial Sequence
249 <220> FEATURE:
250 <223> OTHER INFORMATION: synthetic DNA
252 <400> SEQUENCE: 9
253 tcatggttct tcatgatgat gatgatgatg aactggaaag tggtcgctga catccag 57
256 <210> SEQ ID NO: 10
257 <211> LENGTH: 22
258 <212> TYPE: DNA
259 <213> ORGANISM: Artificial Sequence
261 <220> FEATURE:
262 <223> OTHER INFORMATION: synthetic DNA
264 <400> SEQUENCE: 10
265 ggccacgcgt cgactagtagtt 22

RAW SEQUENCE LISTING ERROR SUMMARY DATE: 03/25/2003
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Please Note:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:5; Xaa Pos. 4

Invalid Line Length:

The rules require that a line not exceed 72 characters in length. This includes spaces.

Seq#:1; Line(s) 7

VERIFICATION SUMMARY

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L:31 M:258 W: Mandatory Feature missing, <223> Blank for SEQ#:1, Line#:29

L:200 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:5 after pos.:0